

IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/829,504A

DATE: 08/04/2004 TIME: 09:16:05

Input Set : A:\23239-558A.Substitute US.ST25.txt

Output Set: N:\CRF4\08042004\J829504A.raw

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3 <110> APPLICANT: Epstein, David
             Grate, Dilara
      4
      5
             Stanton, Martin
      6
              Diener, John L.
      7
              Wilson, Charles
              McCauley, Thomas
              DeSouza, Errol
     11 <120> TITLE OF INVENTION: Stabilized Aptamers to Platelet Derived Growth Factor and
Their
              Use as Oncology Therapeutics
     14 <130> FILE REFERENCE: 23239-558A
     16 <140> CURRENT APPLICATION NUMBER: 10/829,504A
     17 <141> CURRENT FILING DATE: 2004-04-21
     19 <150> PRIOR APPLICATION NUMBER: 10/762915
     20 <151> PRIOR FILING DATE: 2004-01-21
     22 <150> PRIOR APPLICATION NUMBER: 60/441357
     23 <151> PRIOR FILING DATE: 2003-01-21
     25 <150> PRIOR APPLICATION NUMBER: 60/463095
     26 <151> PRIOR FILING DATE: 2003-04-15
     28 <150> PRIOR APPLICATION NUMBER: 60/464179
     29 <151> PRIOR FILING DATE: 2003-04-21
     31 <150> PRIOR APPLICATION NUMBER: 60/465055
     32 <151> PRIOR FILING DATE: 2003-04-23
     34 <150> PRIOR APPLICATION NUMBER: 60/469628
     35 <151> PRIOR FILING DATE: 2003-05-08
     37 <150> PRIOR APPLICATION NUMBER: 60/474680
     38 <151> PRIOR FILING DATE: 2003-05-29
     40 <150> PRIOR APPLICATION NUMBER: 60/491019
     41 <151> PRIOR FILING DATE: 2003-07-29
     43 <150> PRIOR APPLICATION NUMBER: 60/512071
     44 <151> PRIOR FILING DATE: 2003-10-17
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     55 <150> PRIOR APPLICATION NUMBER: 60/428,102
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130 <222> LOCATION: (7)..(7) 131 <223> OTHER INFORMATION: gm.

134 <221> NAME/KEY: modified base

133 <220> FEATURE:

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DATE: 08/04/2004

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Input Set : A:\23239-558A.Substitute US.ST25.txt

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- 135 <222> LOCATION: (10)..(10)
- 136 <223> OTHER INFORMATION: 2' Fluoro-Uracil
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- 140 <222> LOCATION: (11)..(11)
- 141 <223> OTHER INFORMATION: 2' Fluoro-Cytosine
- 143 <220> FEATURE:
- 144 <221> NAME/KEY: modified base
- 145 <222> LOCATION: (12)..(12)
- 146 <223> OTHER INFORMATION: 2'-O-methyl-Adenosine
- 148 <400> SEQUENCE: 2
- 149 cgtagagcau ca
- 152 <210> SEQ ID NO: 3
- 153 <211> LENGTH: 9
- 154 <212> TYPE: DNA
- 155 <213> ORGANISM: Artificial
- 157 <220> FEATURE:
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- 161 <220> FEATURE:
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- 163 <222> LOCATION: (5)..(6)
- 164 <223> OTHER INFORMATION: 2'-fluoro-Cytosine
- 166 <220> FEATURE:
- 167 <221> NAME/KEY: modified_base
- 168 <222> LOCATION: (7)..(7)
- 169 <223> OTHER INFORMATION: 2'-fluoro-Uracil
- 171 <220> FEATURE:
- 172 <221> NAME/KEY: modified base
- 173 <222> LOCATION: (8)..(8)
- 174 <223> OTHER INFORMATION: gm
- 176 <400> SEQUENCE: 3
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- 180 <210> SEQ ID NO: 4
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- 182 <212> TYPE: DNA
- 183 <213> ORGANISM: Artificial
- 185 <220> FEATURE:
- 186 <223> OTHER INFORMATION: aptamer
- 189 <220> FEATURE:
- 190 <221> NAME/KEY: modified_base
- 191 <222> LOCATION: (4)..(4)
- 192 <223> OTHER INFORMATION: 2'-fluoro-Cytosine
- 194 <220> FEATURE:
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- 196 <222> LOCATION: (4)..(4)
- 197 <223> OTHER INFORMATION: 2'-fluoro-cytosine
- 199 <220> FEATURE:
- 200 <221> NAME/KEY: modified base /
- 201 <222> LOCATION: (5)..(5)

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Input Set : A:\23239-558A.Substitute US.ST25.txt

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210 <221> NAME/KEY: modified base
211 <222> LOCATION: (8)..(8)
212 <223> OTHER INFORMATION: 2'-fluoro-Cytosine
214 <220> FEATURE:
215 <221> NAME/KEY: modified_base
216 <222> LOCATION: (9)..(9)
217 <223> OTHER INFORMATION: gm
219 <400> SEQUENCE: 4
220 cagcgwadg/
223 <210> SEQ ID NO: 5
224 <211> LENGTH: 12
225 <212> TYPE: DNA
226 <213> ORGANISM: Artificial
228 <220> FEATURE:
229 <223> OTHER INFORMATION: aptamer
232 <220> FEATURE:
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234 <222> LOCATION: (7)..(7)
235 <223> OTHER INFORMATION: gm
237 <220> FEATURE:
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239 <222> LOCATION: (10)..(10)
240 <223> OTHER INFORMATION: 2'-fluoro-Uracil
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243 <221> NAME/KEY: modified base
244 <222> LOCATION: (11)..(11)
245 <223> OTHER INFORMATION: 2'-fluoro-Cytosine
247 <220> FEATURE:
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250 <223> OTHER INFORMATION: 2'-methyl-Adenosine
252 <220> FEATURE:
253 <221> NAME/KEY: modified base
254 <222> LOCATION: (12)..(12)
255 <223> OTHER INFORMATION: 2'-O-methyl-adenosine
257 <400> SEQUENCE: 5
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264 <213> ORGANISM: Artificial
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267 <223> OTHER INFORMATION: aptamer

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Input Set : A:\23239-558A.Substitute US.ST25.txt

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271 <221> NAME/KEY: modified_base
272 <222> LOCATION: (6)..(6)
273 <223> OTHER INFORMATION: 2'-fluoro-Cytosine
275 <220> FEATURE:
276 <221> NAME/KEY: modified_base
277 <222> LOCATION: (7)...(7)
278 <223> OTHER INFORMATION: 2'-fluoro-Uracil
280 <220> FEATURE:
281 <221> NAME/KEY: modified_base/
282 <222> LOCATION: (8)..(8)
283 <223> OTHER INFORMATION: gm
285 <400> SEQUENCE: 6
286 tgáagcúgt
289 <210> SEQ ID NO: 7
290 <211> LENGTH: 23
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292 <213> ORGANISM: Artificial
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295 <223> OTHER INFORMATION: aptamer
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301 <223> OTHER INFORMATION: 2'-O-methyl-adenosine
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304 <221> NAME/KEY: modified_base
305 <222> LOCATION: (2)..(2)
306 <223> OTHER INFORMATION: 2'-O-methyl-uracil
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309 <221> NAME/KEY: modified base
310 <222> LOCATION: (3)..(3)
311 <223> OTHER INFORMATION: gm
313 <220> FEATURE:
314 <221> NAME/KEY: modified_base
315 <222> LOCATION: (4)..(4)
316 <223> OTHER INFORMATION: cm
318 <220> FEATURE:
319 <221> NAME/KEY: modified_base
320 <222> LOCATION: (5)..(5)
321 <223> OTHER INFORMATION: 2'-O-methyl-adenosine
323 <220> FEATURE:
324 <221> NAME/KEY: modified base
325 <222> LOCATION: (6)..(6)
326 <223> OTHER INFORMATION: gm
328 <220> FEATURE:
329 <221> NAME/KEY: modified base
330 <222> LOCATION: (7)..(9)
331 <223> OTHER INFORMATION: 2'-O-methyl-uracil
333 <220> FEATURE:
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/829,504A

DATE: 08/04/2004 TIME: 09:16:06

Input Set : A:\23239-558A.Substitute US.ST25.txt

Output Set: N:\CRF4\08042004\J829504A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:50; N Pos. 25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43,44 Seq#:50; N Pos. 45,46,47,48,49,50,51,52,53,54 Seq#:51; N Pos. 24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43 Seq#:51; N Pos. 44,45,46,47,48,49,50,51,52,53 Seq#:52; N Pos. 24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43 Seq#:52; N Pos. 44,45,46,47,48,49,50,51,52,53 Seq#:91; N Pos. 25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43,44 Seq#:91; N Pos. 45,46,47,48,49,50,51,52,53,54,55,56,57,58,59,60,61,62,63,64

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27 Seq#:28,29,30,31,32,33,34,35,36,37,50,51,52,53,54,55,56,57,58,59,60,61,62,63 Seq#:64,65,66,67,68,69,70,71,72,73,74,75,76,77,78,79,80,81,82,83,84,85,86,87 Seq#:88,89,90,91,92,93,94,95,96,97,98

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/829,504A

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Input Set: A:\23239-558A.Substitute US.ST25.txt
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L:1007 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (38) SEQUENCE:
L:1012 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (39) SEQUENCE:
L:1017 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (40) SEQUENCE:
L:1022 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (41) SEQUENCE:
L:1027 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (42) SEQUENCE:
L:1032 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (43) SEQUENCE:
L:1037 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (44) SEQUENCE:
L:1042 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (45) SEQUENCE:
L:1047 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (46) SEQUENCE:
L:1052 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (47) SEQUENCE:
L:1057 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (48) SEQUENCE:
L:1062 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (49) SEQUENCE:
L:1079 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 after pos.:0
L:1099 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:0
L:1119 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52 after pos.:0
L:2110 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:91 after pos.:0
L:2112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:91 after pos.:60
```